

I. AMENDMENT

In the Specification:

At page 21, lines 23-28 and page 22, lines 1-2, please replace the paragraph with following:

FIG. 11A, FIG. 11B, FIG. 11C, and FIG. 11D, FIG. 11E, FIG. 11F, FIG. 11G, FIG. 11H, and FIG 11I. DNA (SEQ ID NO:34) and predicted amino acid sequences (SEQ ID NO:35) for UC332 (KA000262) taken from Nagase *et al.* (1996) (GB:D87451). Indicated are a C₃HC₄ zinc RING finger motif (shaded and underlined with conserved cysteines and histidines in bold) located between amino acids 175 and 216, a nuclear localization signal (underlined), a putative leucine zipper sequence (shaded area with repeating leucines and isoleucines in bold), and a PEST sequence flanked by basic stretches of amino acids (underlined) located between amino acid positions 684-736.